|          | CIPErrors Corrected by the STIC Sy GENTED 10 4/12 /200  |
|----------|---|
| Sorbil   | Number: 09/723,000 ENTEREAD and by: 4/2/2001 Changed a file from non-ASCII to ASCII   |
| $\Box$ . | TECH CENTED 4000 to accurate  |
|          | Changed the margins in cases where the sequence text was TECH CENTER 1600/2968 line.  |
|          | Edited a format error in the Current Application Data section, specifically:  |
|          | Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other   |
|          | Added the mandatory heading and subheadings for "Current Application Data".   |
|          | Edited the Number of Sequences field. The applicant spelled out a number instead of using an integer.   |
|          | Changed the spelling of a mandatory field (the headings or subheadings), specifically:  |
|          | Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were edited were:   |
|          | Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited:  |
|          | Corrocted subhoading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  |
|          | Inserted colons after headings/subheadings. Headings edited included:   |
|          | Doloted extra, invalid, headings used by an applicant, specifically:  |
|          | Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of file page numbers throughout text; other invalid text, such as  |
|          | Insorted mandatory headings, specifically:  |
|          | Corrected an obvious error in the response, specifically:   |
|          | Edited identifiers where upper case is used but lower case is required, or vice versa.  |
|          | Corrected an error in the Number of Sequences field, specifically:  |
|          | A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted   |
|          | Deloted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected:   |
|          | Other:  |
|          | James Committee of the |
|          | · · · · · · · · · · · · · · · · · · ·   |

Action. DO NOT send a copy of this form.

1645

RAW SEQUENCE LISTING

DATE: 04/12/2001 PATENT APPLICATION: US/09/723,000 TIME: 16:24:50

Input Set : A:\07039-170002.txt

Output Set: N:\CRF3\04122001\I723000.raw

Does Not Comply Corrected Diskette Needed

```
4 <110> APPLICANT: Goronzy, Jorg J.
        Weyand, Cornelia M.
 7 <120> TITLE OF INVENTION: METHODS AND MATERIALS FOR EVALUATING
        RHEUMATOID ARTHRITIS
10 <130> FILE REFERENCE: 07039-170002
12 <140> CURRENT APPLICATION NUMBER: US 09/723,000.
13 <141> CURRENT FILING DATE: 2000-11-27
15 <150> PRIOR APPLICATION NUMBER: US 09/387,467
16 <151> PRIOR FILING DATE: 1999-09-01
18 <150> PRIOR APPLICATION NUMBER: US 60/098,718
19 <151> PRIOR FILING DATE: 1998-09-01
```

## ERRORED SEQUENCES

```
245 <210> SEQ ID NO: 21
     246 <211> LENGTH: 24
     247 <212> TYPE: DNA
     248 <213> ORGANISM: Artificial Sequence
     250 <220> FEATURE:
     251 <223> OTHER INFORMATION: biotinylated probe
     253 <400> SEQUENCE: 21
     254 aatggcgtgg agctgagaga taac
E--> 258 1
```

23 <170> SOFTWARE: FastSEQ for Windows Version 4.0

21 <160> NUMBER OF SEQ ID NOS: 21

24

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/723,000

DATE: 04/12/2001

TIME: 16:24:51

Input Set : A:\07039-170002.txt
Output Set: N:\CRF3\04122001\I723000.raw

L:258 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:24 SEQ:21